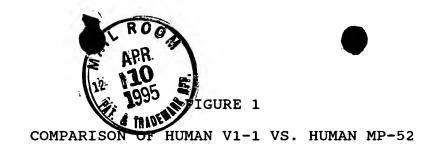
U 8/217780



V1-1 1 MP52 1	AGC GCT	CGC	TGC TGC	AGC AGT	CGC CGG	AAG AAG	CCG - GCA	TTG CTG	CAC	GTG GTC	Asp GAC AAC Asn	TTC	AAG AAG	GAG GAC	CTC ATG
16 46 46 16	GGC GGC	TGG TGG	GAC	GAC	TGG TGG	ATC ATC	ATC ATC	GCG GCA	CCG	CTG CTT	Asp GAC GAG Glu	TAC	GAG GAG	GCG GCT	TAC TTC
31 91 91 31	CAC	TGC	GAG GAG	GGC GGG	CTT CTG	TGC	GAC GAG	TTC TTC	CCT CCA	TTG TTG	Arg CGT CGC Arg	TCG TCC	CAC CAC	CTC CTG	GAG GAG
46 121 121 46	CCC	ACC ACG	AAC AAT	CAT CAT	GCC GCA	ATC GTC	ATT ATC	CAG CAG	ACG ACC	CTG CTG	Leu CTC - ATG Met	AAC AAC	TCC	ATG ATG	GCA GAC
61 181 181 61	CCA CCC	GAC GAG	GCG - TCC	GCG - ACA	CCG CCA	GCC CCC	TCC ACC	TGC TGC	TGT TGT	GTG GTG	Pro CCA !! CCC Pro	GCG ACG	CGC CGG	CTC CTG	AGC AGT
76 226 226 76	CCC	ATC ATC	AGC AGC	ATC ATC	CTC CTC	TAC TTC	ATC ATT	GAC GAC	GCC TCT	GCC GCC	Asn AAC !!! AAC Asn	AAC AAC	GTT GTG	GTC GTG	TAC
91 271 271 91	AAG AAG	CAA CAG	TAC	GAG GAG	GAC GAC	ATG ATG	GTC GTC	GTG GTG	GAG GAG	GCC TCG	Cys TGC TGT Cys	GGC GGC	TGC TGC	AGG AGG	

Homology at the nucleotide level: 249/312 = 79.8%Homology at the amino acid level: 84/104 = 80.8%